=> d his

(FILE 'HOME' ENTERED AT 10:30:43 ON 14 MAR 2008)

FILE 'REGISTRY' ENTERED AT 10:30:50 ON 14 MAR 2008

L1 STRUCTURE UPLOADED

L2 0 S L1

L3 17 S L1 FULL

FILE 'CAPLUS' ENTERED AT 10:31:47 ON 14 MAR 2008

L4 19 S L3

L5 19 S L4 AND PY<=2006

FILE 'REGISTRY' ENTERED AT 10:45:37 ON 14 MAR 2008

L6 STRUCTURE UPLOADED

L7 0 S L6

L8 14 S L6 FULL

FILE 'REGISTRY' ENTERED AT 10:46:50 ON 14 MAR 2008

FILE 'CAPLUS' ENTERED AT 10:46:53 ON 14 MAR 2008

L9 10 S L8

FILE 'REGISTRY' ENTERED AT 10:54:53 ON 14 MAR 2008

L10 STRUCTURE UPLOADED

L11 6 S L10

L12 125 S L10 FULL

FILE 'CAPLUS' ENTERED AT 10:56:13 ON 14 MAR 2008

L13 19 S L12

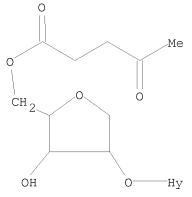
=> d stat que

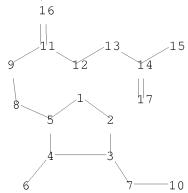
L10 STR

Structure attributes must be viewed using STN Express query preparation. L12 $\,$ 125 SEA FILE=REGISTRY SSS FUL L10 $\,$

L13 19 SEA FILE=CAPLUS ABB=ON PLU=ON L12 =>

Uploading C:\Program Files\Stnexp\Queries\10597445-14Mar08-1.str





chain nodes :

6 7 8 9 10 11 12 13 14 15 16 17

ring nodes :
1 2 3 4 5
chain bonds :

3-7 4-6 5-8 7-10 8-9 9-11 11-12 11-16 12-13 13-14 14-15 14-17

ring bonds :

1-2 1-5 2-3 3-4 4-5

exact/norm bonds :

 $1-2 \quad 1-5 \quad 2-3 \quad 3-4 \quad 3-7 \quad 4-5 \quad 4-6 \quad 7-10 \quad 9-11 \quad 11-16 \quad 14-17$

exact bonds :

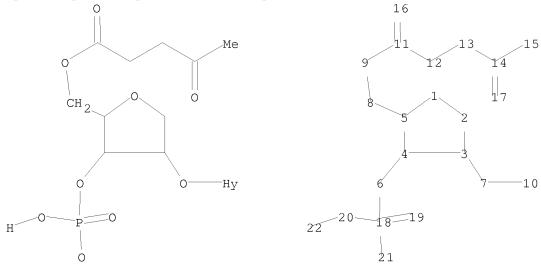
5-8 8-9 11-12 12-13 13-14 14-15

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS 7:CLASS 8:CLASS 9:CLASS 10:Atom 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS

=>

Uploading C:\Program Files\Stnexp\Queries\10597445-14Mar08-.str



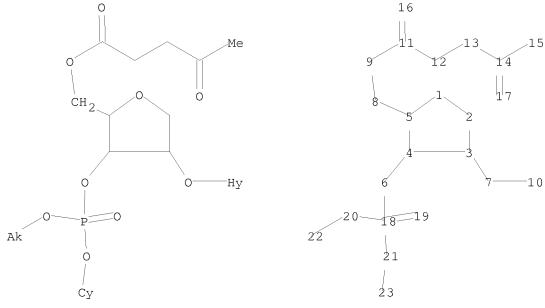
chain nodes : 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 ring nodes : 1 2 3 4 5 chain bonds : 3-7 4-6 5-8 6-18 7-10 8-9 9-11 11-12 11-16 12-13 13-14 14-15 14-1718-19 18-20 18-21 20-22 ring bonds : 1-2 1-5 2-3 3-4 4-5 exact/norm bonds : $1-2 \quad 1-5 \quad 2-3 \quad 3-4 \quad 3-7 \quad 4-5 \quad 4-6 \quad 6-18 \quad 7-10 \quad 9-11 \quad 11-16 \quad 14-17 \quad 18-21$ exact bonds : 5-8 8-9 11-12 12-13 13-14 14-15 20-22 normalized bonds : 18-19 18-20

Match level:

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS 7:CLASS 8:CLASS 9:CLASS 10:Atom 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS 20:CLASS 21:CLASS 22:CLASS

=>

Uploading C:\Program Files\Stnexp\Queries\10597445-14Mar08-3.str



chain nodes :
6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23
ring nodes :
1 2 3 4 5

chain bonds : 3-7 4-6 5-8 6-18 7-10 8-9 9-11 11-12 11-16 12-13 13-14 14-15 14-17 18-19

18-20 18-21 20-22 21-23

ring bonds :

1-2 1-5 2-3 3-4 4-5

exact/norm bonds :

 $1-2 \quad 1-5 \quad 2-3 \quad 3-4 \quad 3-7 \quad 4-5 \quad 4-6 \quad 6-18 \quad 7-10 \quad 9-11 \quad 11-16 \quad 14-17 \quad 18-19 \quad 18-20$

18-21 20-22 21-23

exact bonds :

5-8 8-9 11-12 12-13 13-14 14-15

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS 7:CLASS 8:CLASS 9:CLASS 10:Atom 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS

21:CLASS 22:CLASS 23:Atom

=> d his

(FILE 'HOME' ENTERED AT 07:52:07 ON 17 MAR 2008)

FILE 'REGISTRY' ENTERED AT 07:52:22 ON 17 MAR 2008 E LIPASE/CN

FILE 'CAPLUS' ENTERED AT 07:52:53 ON 17 MAR 2008

L3 35400 S L2 5363 S LEVULIN? L4104 S LEVULINOYL L5 1327 S LEVULINATE L6 L7 4043 S LEVULINIC ACID L8 4884 S L5 OR L6 OR L7 1 S L2 (S) L8 L9 1 S L2 (L) L8 L10

16 S L2 AND L8

=> d stat que

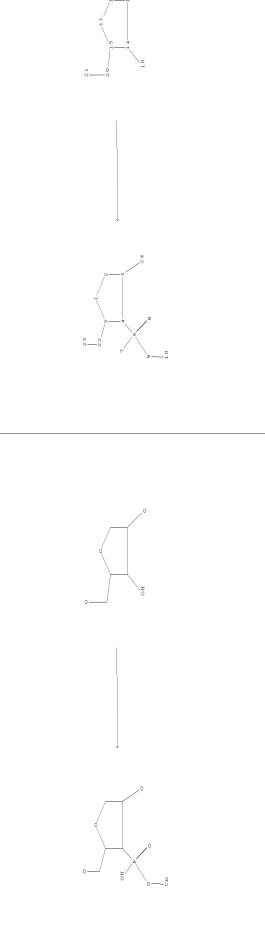
L11

L2 1 SEA FILE=REGISTRY ABB=ON PLU=ON LIPASE/CN
L5 104 SEA FILE=CAPLUS ABB=ON PLU=ON LEVULINOYL
L6 1327 SEA FILE=CAPLUS ABB=ON PLU=ON LEVULINATE
L7 4043 SEA FILE=CAPLUS ABB=ON PLU=ON LEVULINIC ACID
L8 4884 SEA FILE=CAPLUS ABB=ON PLU=ON L5 OR L6 OR L7
L11 16 SEA FILE=CAPLUS ABB=ON PLU=ON L2 AND L8

```
FILE 'CASREACT' ENTERED AT 11:11:30 ON 20 MAR 2008
               STRUCTURE UPLOADED
L1
L2
              0 S L1
L3
              0 S L2 FULL
L4
              0 S L1 FULL
L5
                STRUCTURE UPLOADED
L6
              0 S L5
L7
              0 S L5 FULL
     FILE 'STNGUIDE' ENTERED AT 11:14:52 ON 20 MAR 2008
     FILE 'REGISTRY' ENTERED AT 11:16:14 ON 20 MAR 2008
               E 2-CHLOROPHENYLPHOSPHOROTRIAZOLIDE
                E 2-CHLOROPHENYLPHOSPHOROTRIAZOLIDE/CN
                E PHOSPHOROTRIAZOLIDE/CN
           1668 S CHLORO (XA) PHOSPHORO (XA) PHENYL (XA) TRI
Γ8
              O S CHLORO (XA) PHOSPHORO (XA) PHENYL (XA) TRIAZOLIDE
L9
L10
            197 S CHLORO (XA) PHOSPHORO (XA) PHENYL (XA) TRIAZO?
            197 S CHLORO (XA) PHOSPHORO (XA) PHENYL (XA) TRIAZO? (XA) 2
L11
     FILE 'CAPLUS' ENTERED AT 11:19:04 ON 20 MAR 2008
                E US2007172925/PN
L12
              1 S E3
     FILE 'REGISTRY' ENTERED AT 11:20:45 ON 20 MAR 2008
L13
              1 S 861817-06-3/RN
                SET NOTICE 1 DISPLAY
                SET NOTICE LOGIN DISPLAY
     FILE 'CAPLUS' ENTERED AT 11:21:40 ON 20 MAR 2008
                SEL L12 RN
     FILE 'REGISTRY' ENTERED AT 11:21:50 ON 20 MAR 2008
L14
            85 S E1-E85
L15
             3 S L14 AND PHOSPH?
     FILE 'CASREACT' ENTERED AT 11:22:47 ON 20 MAR 2008
            48 S L15
L16
     FILE 'REGISTRY' ENTERED AT 11:26:25 ON 20 MAR 2008
L17
             1 S L14 AND TRICHLORO?
     FILE 'CASREACT' ENTERED AT 11:27:09 ON 20 MAR 2008
            264 S L17
L18
             48 S L18 AND PHOSPH?
L19
     FILE 'CASREACT' ENTERED AT 11:40:02 ON 20 MAR 2008
             9 S L19 AND PHOSPHITE
L20
     FILE 'STNGUIDE' ENTERED AT 11:50:16 ON 20 MAR 2008
=> d stat que
             85 SEA FILE=REGISTRY ABB=ON PLU=ON (100-07-2/BI OR 108-24-7/BI
L14
                OR 109-78-4/BI OR 115-20-8/BI OR 118-00-3/BI OR 1191-99-7/BI
                OR 120187-92-0/BI OR 120187-94-2/BI OR 138494-30-1/BI OR
                21090-30-2/BI OR 40608-06-8/BI OR 40615-36-9/BI OR 40615-37-0/B
                I OR 40615-39-2/BI OR 64350-24-9/BI OR 65-46-3/BI OR 69304-37-6
                /BI OR 72351-28-1/BI OR 75-77-4/BI OR 79-30-1/BI OR 80817-46-5/
                BI OR 85393-37-9/BI OR 861816-91-3/BI OR 861816-92-4/BI OR
```

861816-93-5/BI OR 861816-94-6/BI OR 861816-95-7/BI OR 861816-97 -9/BI OR 861816-98-0/BI OR 861816-99-1/BI OR 861817-01-8/BI OR 861817-03-0/BI OR 861817-04-1/BI OR 861817-05-2/BI OR 861817-06 -3/BI OR 861817-07-4/BI OR 861817-09-6/BI OR 861817-10-9/BI OR 861817-11-0/BI OR 861842-82-2/BI OR 861842-83-3/BI OR 861842-85 -5/BI OR 861842-86-6/BI OR 861842-87-7/BI OR 861842-88-8/BI OR 861842-89-9/BI OR 861842-90-2/BI OR 861842-91-3/BI OR 861842-92 -4/BI OR 861842-93-5/BI OR 861842-95-7/BI OR 861842-96-8/BI OR 861842-97-9/BI OR 861842-98-0/BI OR 861842-99-1/BI OR 861843-00 -7/BI OR 861843-01-8/BI OR 861843-02-9/BI OR 861843-03-0/BI OR 861843-04-1/BI OR 861843-05-2/BI OR 861843-06-3/BI OR 861843-07 -4/BI OR 861843-08-5/BI OR 861843-09-6/BI OR 861843-10-9/BI OR 861843-11-0/BI OR 861843-12-1/BI OR 861843-13-2/BI OR 862202-73 -1/BI OR 862224-10-0/BI OR 862224-11-1/BI OR 862224-12-2/BI OR 862224-13-3/BI OR 862224-14-4/BI OR 862224-15-5/BI OR 862224-16 -6/BI OR 862224-17-7/BI OR 862224-18-8/BI OR 862259-89-0/BI OR 86327-76-6/BI OR 87865-78-9/BI OR 9001-62-1/BI OR 90865-73-9/BI OR 90865-74-0/BI)

L17 1 SEA FILE=REGISTRY ABB=ON PLU=	=ON LI4 AND IRICHLORO?
L18 264 SEA FILE=CASREACT ABB=ON PLU=	ON L17
L19 48 SEA FILE=CASREACT ABB=ON PLU=	ON L18 AND PHOSPH?
L20 9 SEA FILE=CASREACT ABB=ON PLU=	ON L19 AND PHOSPHITE



-15 14 22 - 213-19 20-21 -14 13 15-20 14 - 14 14-16 13 15 -13 13-19 Ω 4 12 -12 Ŋ 9-10 H $_{\rm U}^{\rm U}$ Ŋ 9 - 12 И Ц 6-9 14 Ŋ 1 4 0 13 8 0 Ŋ 9 12 Н 16 0 11 $^{\circ}$ 1 2 2 10 Ŋ ring bonds: 1-2 1-5 2-3 exact/norm bonds 22-23 1-5 2-3 σ chain nodes < 7 8 ring nodes 1 2 3 chain bonds

15-2 exact bonds: 4-6 5-22 9-10 normalized bonds: 6-7 6-8

3:Atom 4:Atom 5:Atom 6:CLASS 7:CLASS 8:CLASS 14:Atom 15:Atom 16:CLASS 19:CLASS 20:CLASS : 2:Atom 3 13:Atom Match level:
1:Atom 2
12:Atom
24:CLASS
fragments ass

11:Atom 23:CLASS

10:Atom 22:CLASS

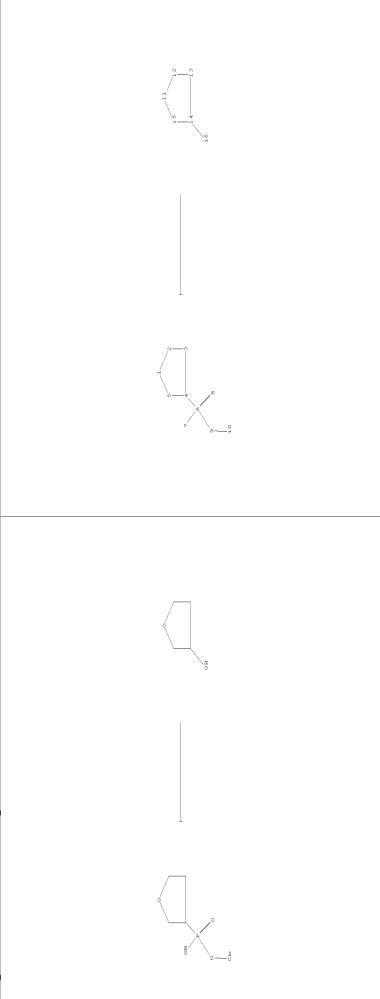
9:CLASS 21:CLASS

20-21

14-16

role:

node mappings: 4:14 4:14



11-12 6 - 9 Ŋ 4 3 - 4 ring bonds:

1-2 1-5 2-3

exact/norm bonds:

1-2 1-5 2-3

exact bonds:

4-6 9-10

normalized bonds:

6-7 6-8

9:CLASS 8:CLASS 7:CLASS Match level:
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS
12:Atom 13:Atom 14:Atom 15:Atom 16:CLASS
reaction site bonds:
4-6:XC 14-16:XC

11:Atom

10:Atom

14-16

14 - 15

-14

12

11-15

14-15

13-14

12 - 13

11-15